

Page 1

OM nucleic - nucleic search, using sw model GenCore version 5.1.7 Copyright (c) 1993 - 2006 Biocceleration Ltd.

a O

Run on:

Pebruary 10, 2006, 05:26:42 ; Search time 4815 Seconds (without alignments)
10270.779 Million cell updates/sec

Title: Perfect score: Sequence:

Scoring table: IDENTITY NUC Gapop 10.0 , Gapext 1.0

5883141 seqs, 28421725653 residues

Searched:

U8-10-600-997-7 870 1 atgaagacattgcctgccat.....ccatatgtgtgaggagttaa 870

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Total number of hits satisfying chosen parameters:

11766282

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : GenEmbl:*
1: gb ba:*
2: gb in:*
3: gb on:*
5: gb oc:*
6: gb pa:*
7: gb pa:*
9: gb pa:*
9: gb re:*
10: gb sts:*
11: gb sts:*
11: gb sts:*
12: gb n:*
5: gb pa:*
9: gb ro:*
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9: gb ro:*
9: gb pa:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Regult	ъ	N	w	۱ 4	1	U	6	7	0		10	H	12	13	14	15	16	17	18
Score	870	857.2	857.2	954	0.04	846	764.2	704.2	527.6	520.2	466.2	437.6	425	425	402.8	402.8	402.8	388	378
Query	100.0	98.5	98.5	5	7.86	97.2	87.8	80.9	60.6		53.6	50.3	48.9	48.9	46.3	46.3	46.3	44.6	43.4
Length	870	870	990	1066	1066	3002	777	717	534	726	471	444	769	769	1903	1903	1903	396	1940
80		O)	J.	n (σ	8	σ	0	Ġ	σ	o	o	σ	σ	σ	σ	o	o	o
ID	AY293286	C0947366	C0947428		CS105855	AK131204	CQ947365	CQ947364	CQ947360	BD060983	CQ947358	CQ947359	C8035517	CS044469	BD131124	CS132603	AR339778	CQ947363	BD131144
Description	АУ293286 Н				CS105855 S	AK131204 H	-	CQ947364 S			CQ947358 S	CQ947359 S	CS035517 ·8	CS044469 S					
ង	Homo sapi	Section Car	odience od	actrance	Sequence	Homo sapi	Sequence	Sequence	Sequence	Secreted	Sequence	Sequence	Sequence	Sequence	45 human	Sequence	Sequence	Sequence	45 human

149.2	156.4	157.2	157.2	198.4	199.4	233	233	236.8	254.2	258.8	264.6	•	267.6	267.6	267.6	275.6	276.6	276.6	285.2	315.8	315.8	315.8	326.6	376.2	378	378
17.1	18.0	18.1	18.1	22.8	22.9	26.8	26.8	27.2	29.2	29.7	30.4	30.4	30.8	30.8	30.8	31.7	31.8	31.8	32.8	36.3	36.3	36.3	37.5	43.2	43.4	43.4
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_	7405														CQ947408 Sequence		AX780935 Sequence	AX779565 Sequence	Rattu		AC024131 Homo sapi			CQ947357 Sequence	_	CS132623 Sequence

ALIGNMENTS

	CDS	gene	source	FEATURES	JOURNAL	AUTHORS	JOURNAL REFERENCE		TITLE	REFERENCE AUTHORS		KEYWORDS SOURCE ORGANISM	ACCESSION VERSION	RESULT 1 AY293286 LOCUS DEFINITION
/gene="BTLA" /note="immunoregulatory inhibitory receptor; similar to /note="immunoregulatory inhibitory receptor; similar to CTLA-4 and PD-1" /codon_start=1 /product="B and T lymphocyte attenuator" /product="B and T lymphocyte attenuator" /protein_id="ApP44003.1" /db_xref="GI:31880027"	/gene="BTLA" 1870	/organism="Homo sapiens" /mol_type="mRNA" /db_xref="taxon:9606" /ceIl line="Ramos" 1. 870		3110, USA fiers		Murphy, K.M., Watanabe, N., Yang, J. and Murphy, T.L.	Nat. Immunol. (2003) In press 2 (bases 1 to 870)	CTLA-4 and PD-1	, Zimmerman, N., Sim, J., Zang, X., Allison, J.P. and Murphy, K.M.	<pre>1 (bases 1 to 870) Watanabe, N., Gavrieli, M., Sedy, J.R., Yang, J., Fallarino, F.,</pre>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae; Homo.	Homo sapiens (human) Homo sapiens	CGB. AY293286 AY293286.1 GI:31880026	AY293286 870 bp mRNA linear PRI 18-JUN-2003 Homo sapiens B and T lymphocyte attenuator (BTLA) mRNA, complete

/tranelation="MKTLPAMLGTGKLFWVFFLIPYLDIWNIHGKESCDVQLYIKRQS EHSILAGDPFELECPVKYCANRPHVTWCKLNGTTCVKLEDRQYSWKEEKNISFFILHF EPMLFNDNGSYRCSENFOSNLISSTYADTVKGASEFSKDEVASRPWLLYSLLFLGGLFLLITTWFCLFCCLRRHQKGNSTSTYADTVKGASEFSKDEVASRPWLLYSLLFLGGLFLLITTWFCLFCCLRRHQKGNSLSDTARBINLVDAHLKSEQTEASTRQNSQV LLSEAGIYDNDFDLCFRWQEGSEVCSNFCLEENKFGIVYASLNHSVIGLNSRLARNVK EAPTEYASICVRS"

8 δ र् ð रु है री 문 र् 밁 S 밁 S 문 8 문 밁 닭 र् 밁 ð 밁 문 र् 문 र् Query Match 100.0%; Score 870; DB 8; I Best Local Similarity 100.0%; Pred. No. 3.6e-224; Matches 870; Conservative 0; Mismatches 0; 109 541 481 301 241 841 781 781 721 721 199 199 109 541 481 421 421 361 361 301 241 181 181 121 121 AAGAGACAATCTGAACACTCCATCTTAGCAGGAGATCCCTTTGAACTAGAATGCCCTGTG 180 61 13 CCATATCTGGACATCTGGAACATCCATGGGAAAGAATCATGTGATGTACAGCTTTATATA 120 GAAGTITGITCTAATCCATGCCTGGAAGAAAACAAACCAGGCATTGITTATGCTTCCCTG 780 GCTCACCTTAAGAGCGAGCAAACAGAAGCAAGCACCAGGCAAAATTCCCCAAGTACTGCTA 660 CACCAAGGAAAGCAAAATGAACTCTCTGACACAGCAGGAAGGGAAATTAATCTGGTTGAT GAACGACCCTCCAAGGACGAAGTGGCAAGCAGACCCTGGCTCCTGTATAGTTTACTTCCT GAACGACCCTCCAAGGACGAAGTGGCAAGCAGACCCTGGCTCCTGTATAGTTTTACTTCCT CATTTTGAACCAATGCTTCCTAATGACAATGGGTCATACCGCTGTTCTGCAAATTTTCAG CATTITGAACCAATGCTTCCTAATGACAATGGGTCATACCGCTGTTCTGCAAATTTTCAG 360 AAACTTGAAGATAGACAAACAAGTTGGAAGGAAGAACATTTCATTTTTCATTCTA AAACTTGAAGATAGACAAACAAGTTGGAAGGAAGAAGAAGAACATTTCATTTTTCATTCTA 300 AAGAGACAATCTGAACACTCCATCTTAGCAGGAGATCCCTTTGAACTAGAATGCCCTGTG CCATATCTGGACATCTGGAACATCCATGGGAAAGAATCATGTGATGTACAGCTTTATATA AACCATTCTGTCATTGGACTGAACTCAAGACTGGCAAGAAATGTAAAAGAAGCACCAACA 840 TCAGAAAGCTGGAATTTATGATAATGACCCTGACCTTTGTTTCAGGATGCAGGAAGGGTCT 720 GCTCACCTTAAGAGCGAGCAAACAGAAGCAAGCACCAGGCAAAATTCCCCAAGTACTGCTA TIGGGGGGATIGCCICTACTCATCACTACCIGGTTCTGCCIGCTGCTGCCTGAGAAGG TIGGGGGGATIGCCTCTACTCATCACTACCTGGTTCTGCCTGTTCTGCTGCCTGAGAAGG GAATATGCATCCATATGTGTGAGGAGTTAA 870 TCAGAAGCTGGAATTTATGATAATGACCCTGACCTTTGTTTCAGGATGCAGGAAGGGTCT 0 Length 870; Indels <u>,</u> Gaps 600 540 540 480 480 300 120 60 60 360 180 600 840 660 720

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GAATATGCATCCATATGTGTGAGGAGTTAA 870

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601 GCTCACCTTAAGAGTGAGCAAACAGAAGCAAGCACCAGGCAAAATTTCCCAAGTACTGCTA 660

RESULT 2	
LOCUS LOCUS	CQ947366 870 bp DNA linear PAT 01-DEC-2004 Sequence 42 from Patent WO2004096976.
ACCESSION VERSION	03
S	Comp apprions (himan)
SIN	hordata: Craniata: Vertebrata: Ruteleostomi
	uarchontoglires; Primates; Catarrhini;
REFERENCE AUTHORS TITLE JOURNAL	1 Raye, J. and Wilkinson, B. Spex compositions and methods of use Spex compositions and methods of use Patent: WO 2004096976-A 42 11-NOV-2004;
FEATURES	Novartis AG (CH); Novartis Pharma GmbH (AT); The Scripps Research Institute (US) Location/Qualifiers
Bource) 1870 /organism="Homo sapiens" /mol_type="unassigned DNA" /db. rref="tayon.9606"
ORIGIN	/ AND PERSON CONTRACTOR CONTRACTO
Query Match Best Local Matches 86	ch 98.5%; Score 857.2; DB 6; Length 870; il Similarity 99.1%; Pred. No. 1e-220; 862; Conservative 0; Mismatches 8; Indels 0; Gaps 0;
Ą	1 ATGAAGACATTGCCTGCCATGCTTGGAACTGGGAAATTATTTTGGGTCTTCTTCTTAATC 60
В	1 ATGAAGACATTGCCTGCCATGCTTGGAACTGGGAAATTATTTTTGGGTCTTCTTAATC 60
γQ	61 CCATATCTGGACATCTGGAACATCCATGGGAAAGAATCATGTGATGTACAGCTTTATATA 120
D _D	61 CCATATCTGGACATCTGGAACATCCATGGGAAAGAATCATGTGATGTACAGCTTTATATAT 120
Q 2	121 AAGAGACAATCTGAACACTCCATCTTAGCAGGAGATCCCTTTGAACTAGAATGCCCTGTG 180
Db 1	121 AAGAGACAATCTGAACACTCCATCTTAGCAGGAGATCCCTTTGAACTAGAATGCCCTGTG 180
Q	181 AAATACTGTGCTAACAGGCCTCATGTGACTTGGTGCAAGCTCAATGGAACAACATGTGTA 240
рb 1	181 AAATACTGTGCTAACAGGCCTCATGTGACTTGGTGCAAGCTCAATGGAACAACATGTGTA 240
Qy 2	241 AAACTTGAAGATAGACAAACTAGTTGGAAGGAAGAGAGAG
р ь 2	241 AAACTIGAAGATAGACAAACAAGTIGGAAGGAAGAAGAACATTICATTI
•	01 CATTITGAACCAATGCTTCCTAATGACAATGGGTCATACCGCTGTTCTGCAAATTITTCAG 36
? 6	2 2
Q to	21 GAACGACCCTCCAAGGACGAAGTGGCAAGCAGACCCTGGCTCCTGTATAGTTTACTTCCT 48
Db 4	421 GAACCACCTCCAAGGACGAAATGGCAAGCAGACCCTGGCTCCTGTATAGTTTACTTCCT 480
Qy	481 TIGGGGGGATIGCCTCIACTCAICACCTACCIGGCTTCTGCCTGTTCTGCCTGAGAAAGG 540
Db	481 TTGGGGGGATTGCCTCTACTCATCACTACCTGTTTCTGCCTGTTCTGCTGCCTGAGAAGG 540
ą	541 CACCAAGGAAAGCAAAATGAACTCTCTGACACAGGAGGAAGGGAAATTAATCTGGTTGAT 600
Db 5	41 CACCAAGGAAAGCAAAATGAACTCTCTGACACAGCAGGAAGGGAAATTAACCTGGTTGAT 6
Q Q	501 GCTCACCTTAAGAGCGAGCAAACAGAAGCAAGCACCAGGCAAAATTCCCCAAGTACTGCTA 660

143

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APPLICANT: HILARY CLARK
APPLICANT: DANIEL L. BATON
APPLICANT: AUSTIN L. GURNEY
APPLICANT: AUSTIN L. GURNEY
APPLICANT: AUSTIN L. GURNEY
APPLICANT: AUSTIN L. GURNEY
APPLICANT: BERND WRANIK
TITLE OF INVENTION: IMMUNE RELATED DISEASES
FILE REFFERENCE: P1996R1-US
FILE REFFERENCE: P1996R1-US
CURRENT APPLICATION NUMBER: US/10/371,341
CURRENT APPLICATION NUMBER: US 60/421,236
PRIOR PILING DATE: 2002-10-25
NUMBER OF SEQ ID NOS: 2
1 SEQ ID NO 1
1 LENGTH: 1066
TYPE: DNA
1 ORGANISM: Homo Sapien
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                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/10371341 Publication No.: US20040091884A1 GENERAL INFORMATION:
                                                               Query Match
Best Local (
                                                Matches
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                                                             Similarity
ATGAAGACATTGCCTGCCATGCTTGGAACTGGGAAATTATTTTGGGTCTTCTTAATC
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98.9%;
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                                           Score 854; DB 7; Pred. No. 2.8e-246; O; Mismatches 10;
                                                                        Length 1066;
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APPLICANT: Genentech, Inc.
APPLICANT: Crowley, Craig
APPLICANT: De Sauvage, Frederic J.
APPLICANT: Baton, Daniel L.
APPLICANT: Ebens, Allen
APPLICANT: Polson, Andrew
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
TITLE OF INVENTION: Compositions ar
                                                                                                                              Sequence 27, Application US/10989826 Publication No. US20050238650A1 GENERAL INFORMATION:
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RESULT 1 US-10-37	222 232 2444 200 200 200 200 200 200 200 200 2
RESULT 1 US-10-371-341-2	739.5 739.5 692.5 692.6 691 673.5 635.5 635.5 572.5 516 516 516 474

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ALIGNMENTS	5 US-10-831-622-99 5 US-10-964-215-99 5 US-10-9131-622-18 5 US-10-964-215-12 5 US-10-964-215-12 5 US-10-964-215-62 5 US-10-964-215-61 5 US-10-964-215-61 5 US-10-964-215-61 5 US-10-964-215-61 6 US-10-964-215-17 7 US-10-964-215-17 8 US-10-831-622-17 9 US-10-964-215-17 6 US-10-964-215-17 7 US-10-964-215-17 8 US-10-918-446-107 9 US-11-002-756-107
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RESULT 2 US-10-600-997-6

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                                                         APPLICANT: Yang, Jianfei
APPLICANT: Yang, Xingxing
APPLICANT: Zang, Xingxing
TITLE OF INVENTION: COMPOSITIONS and Methods for
PILS REFERENCE: A-71608/TAL/DHR
CURRENT APPLICATION NUMBER: US/10/600,997
CURRENT FILLNG DATE: 2003-06-20
PRIOR APPLICATION NUMBER: US 60/390,653
PRIOR APPLICATION NUMBER: US 60/390,653
PRIOR PILING DATE: 2003-01-06
PRIOR PILING DATE: 2003-01-06
NUMBER OF ENGLISH 2003-01-06
NUMBER OF ENGLISH 2003-01-06
NUMBER: DATE: 1005: 56
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APPLICANT: Allison, James
APPLICANT: Murphy, Kenneth
APPLICANT: Watanabe, Norihiko
APPLICANT: Murphy, Theresa
APPLICANT: Murphy, Theresa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 6, Application US/10600997 Publication No. US20040175380A1
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LENGTH: 289
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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APPLICANT: DANIEL L. EATON
APPLICANT: AUSTIN L. GURNEY
APPLICANT: BERND WRANIK
APPLICANT: AUSTIN L. GURNEY
AUSTIN L. G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PILE REPERENCE: P1996R1-US
CURRENT APPLICATION NUMBER: US/10/371,341
CURRENT FILING DATE: 2003-02-19
PRIOR APPLICATION NUMBER: US 60/421,236
PRIOR PILING DATE: 2002-10-25
NUMBER OF SEQ ID NOS: 2
TYPE: PRT
ORGANISM: Homo sapien
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Pred. No. 1.5e-145;
2; Mismatches 4;
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SOFTWARE: PASTSEQ for Windows Version
SEQ ID NO 21
LENGTH: 289
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Publication No. US20040248257A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2004-04-23
PRIOR APPLICATION NUMBER: US 60/467,206
PRIOR FILING DATE: 2003-04-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kaye, Jonathan
APPLICANT: Kilkinson, Beverley
TITLE OF INVENTION: SPEX COMPOSITIONS AND METHODS
FILE REFERENCE: TSRI 810.1
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                     EVCSNPCLBENKPGIVYASLNHSVIGLNSRLARNVKEAPTEYASICVRS 289
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